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IN RE APPLICATION OF

Art Unit: TBA

CADE ET AL.

Examiner: TBA

APPLICATION NO: TBA

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FOR: GENES ENCODING PROTEINS INVOLVED IN THE REGULATION
OF SAR GENE EXPRESSION IN PLANTS

Commissioner for Patents
Washington, D.C. 20231

SUBMISSION OF SEQUENCE LISTING
INCLUDING STATEMENT OF VERIFICATION

Sir:

Applicants hereby provide a Computer Readable Form of the Sequence Listing as well as the Paper Copy thereof. The undersigned states that the Paper Copy and the Computer Readable Form, submitted in accordance with 37 CFR §1.821(c) and (e), respectively, are the same.

Respectfully submitted,



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Date: December 8, 2000

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SEQUENCE LISTING

<110> Cade, Rebecca M
Dietrich, Robert A

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SAR GENE EXPRESSION IN PLANTS

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<150> 60/171,008

<151> 1999-12-15

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aac gga aaa tct gac ggt aac aga ggg aaa ccg tcg acg gaa gtt gtt 157
Asn Gly Lys Ser Asp Gly Asn Arg Gly Lys Pro Ser Thr Glu Val Val
15 20 25 30

- 2 -

cgg acg gta acg gag gaa gag gtg gat gag ttt ttc aag ata tta cgg 205
 Arg Thr Val Thr Glu Glu Val Asp Glu Phe Phe Lys Ile Leu Arg
 35 40 45

aga gta cac gtg gcg aca cga acg gtt gcg aaa gtt aac ggc ggt gtt 253
 Arg Val His Val Ala Thr Arg Thr Val Ala Lys Val Asn Gly Gly Val
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gct gag gga gag tta ccg tct aag aag agg aaa cgg agt cag aat ctt 301
 Ala Glu Gly Glu Leu Pro Ser Lys Lys Arg Lys Arg Ser Gln Asn Leu
 65 70 75

ggg ttg aga aac tcg ttg gat tgt aac ggc gtt cga gac gga gaa ttc 349
 Gly Leu Arg Asn Ser Leu Asp Cys Asn Gly Val Arg Asp Gly Glu Phe
 80 85 90

gat gag att aat cgg gtc ggg tta cag ggt ttg ggt ttg gat ctg aac 397
 Asp Glu Ile Asn Arg Val Gly Leu Gln Gly Leu Gly Leu Asp Leu Asn
 95 100 105 110

tgt aaa ccg gaa cca gac agc gtt agt tta tcg ttg tagacttgta 443
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 35 40 45

His Val Ala Thr Arg Thr Val Ala Lys Val Asn Gly Gly Val Ala Glu
 50 55 60

Gly Glu Leu Pro Ser Lys Lys Arg Lys Arg Ser Gln Asn Leu Gly Leu
 65 70 75 80

Arg Asn Ser Leu Asp Cys Asn Gly Val Arg Asp Gly Glu Phe Asp Glu
 85 90 95

Ile Asn Arg Val Gly Leu Gln Gly Leu Gly Leu Asp Leu Asn Cys Lys

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105

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 ggaaaattca ggtaaaaaga gaaaataaag aatgagagat agagagattt ctatggaaaa 240
 agaaagagag aacatgtagg tgaacaaaat aaagagatat gatgatatat tttatgagag 300
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ctaactattg ccaaaatttc tgtagccgac aaatactatt tgggtccaagg ttattttgtg 540
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ttttccaca catggacttc ctttattcca aaagtcaata aagtgtgacg tcatgatact 660
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tga atg cta ctt atg gac gga gaa aag aag agg aag aga aca gca atc 168
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ggc gcc gga gat cgg agt aag gat gag gta gaa gct act gtg aag gag 216
Gly Ala Gly Asp Arg Ser Lys Asp Glu Val Glu Ala Thr Val Lys Glu
              20              25              30

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gag gag ccg ccg tca gag gcg gag gtt gac gag ttc ttc gcg atc tta 264
Glu Glu Pro Pro Ser Glu Ala Glu Val Asp Glu Phe Phe Ala Ile Leu
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cgg agg atg cat gtg gcg gtg aaa tat ctc cag aga aat gct cag att 312
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 65 70 75

gct gga cgg aag aga gaa cgg gga atc gtg aga aaa ggt gat ttg gac 408
 Ala Gly Arg Lys Arg Glu Arg Gly Ile Val Arg Lys Gly Asp Leu Asp
 80 85 90 95

ctc aac act ctg ccg gac ggc gga gac taa ttaacgcagt ttaagcatag 458
 Leu Asn Thr Leu Pro Asp Gly Gly Asp
 100 105

gttaattaca taaatgcacc cttaattatc gtagattctt aagattgatc tgctgtacag 518

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 Glu Pro Pro Ser Glu Ala Glu Val Asp Glu Phe Phe Ala Ile Leu Arg
 35 40 45
 Arg Met His Val Ala Val Lys Tyr Leu Gln Arg Asn Ala Gln Ile Arg
 50 55 60
 Pro Glu Asn Leu Asn Ala Ser Pro Ala Gly Ala Asn Gly Val Ala Ala
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 85 90 95
 Asn Thr Leu Pro Asp Gly Gly Asp
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<221> CDS

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 cac atg gcc gta aaa tat ctt cag aga aac gct cag att cag ccg gaa 95
 His Met Ala Val Lys Tyr Leu Gln Arg Asn Ala Gln Ile Gln Pro Glu
 20 25 30
 aac gtt aac gct cac ggc agc aag tta acc gca tcg ccg gcc ggt gtt 143
 Asn Val Asn Ala His Gly Ser Lys Leu Thr Ala Ser Pro Ala Gly Val
 35 40 45
 aac gga gat gca act gga cag aag aga gaa cgg gga atc gtg aga aaa 191
 Asn Gly Asp Ala Thr Gly Gln Lys Arg Glu Arg Gly Ile Val Arg Lys
 50 55 60
 ggt gat ttg gac ctc aac act ttg ccg gac tgc gga gac taa 233
 Gly Asp Leu Asp Leu Asn Thr Leu Pro Asp Cys Gly Asp
 65 70 75
 cgcagtttaa gcataggtta attacagaaa tgcaccttta attatcgtag attcttaaga 293
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 20 25 30

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Asp Gly Ser Asp Gly Val Pro Thr Glu Glu Glu Val Glu Glu Phe Phe
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Ala Ile Leu Arg Arg Met Arg Met Ala Val Lys Tyr Phe Asp Asp Lys
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Gly Lys Glu Trp Arg Lys Ala Leu Glu Thr Ala Glu Leu Thr Val Asp
35 40 45

His Arg His Asp Val Val Ala Ala Glu Glu Asp Asp Lys Pro Arg Lys
50 55 60

Lys Gly Gly Glu Val Ile Ile Asn Glu Gly Phe Asp Leu Asn Ala Val
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Ala Pro Glu Ala Ala Glu Gly Gly Gly Ala
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<211> 85

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<213> Nicotiana tabacum

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Thr Ala Val Ser Pro Pro Pro Ser Glu Ala Glu Val Asp Glu Phe Phe
35 40 45

Ala Ile Leu Arg Arg Met His Val Ala Val Arg Tyr Leu Gln Glu Ser
50 55 60

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Gly Gln Lys Arg Val Val Pro Lys Gly Asp Leu Asp Leu Asn Thr Leu
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Pro Gly Asn Gly Asp
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gaa	gaa	aag	atg	gag	aag	ttg	tac	aca	gtg	ctt	aaa	aac	gca	agg	gaa	96
Glu	Glu	Lys	Met	Glu	Lys	Leu	Tyr	Thr	Val	Leu	Lys	Asn	Ala	Arg	Glu	
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atg	cgg	aaa	tat	gtc	aac	agc	tcc	atg	gag	aag	aag	aga	cag	gaa	gaa	144
Met	Arg	Lys	Tyr	Val	Asn	Ser	Ser	Met	Glu	Lys	Lys	Arg	Gln	Glu	Glu	
		35					40					45				

gaa	gaa	aga	gca	agg	gtt	cgt	aga	ttc	cct	tcg	ttt	cag	cca	gaa	gat	192
Glu	Glu	Arg	Ala	Arg	Val	Arg	Arg	Phe	Pro	Ser	Phe	Gln	Pro	Glu	Asp	
	50					55					60					

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Phe	Ile	Phe	Met	Asn	Lys	Ala	Glu	Ala	Asn	Asn	Ile	Glu	Lys	Ala	Ala	
65				70					75					80		

aat	gag	agc	tct	tca	gca	tcc	aac	gag	tat	gat	ggc	tct	aag	gaa	aag	288
Asn	Glu	Ser	Ser	Ser	Ala	Ser	Asn	Glu	Tyr	Asp	Gly	Ser	Lys	Glu	Lys	
			85					90					95			

caa	gaa	gga	tct	gag	act	aac	gtt	tgt	tta	gac	ttg	aat	ctt	tct	ctg	336
Gln	Glu	Gly	Ser	Glu	Thr	Asn	Val	Cys	Leu	Asp	Leu	Asn	Leu	Ser	Leu	
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Met	Arg	Lys	Tyr	Val	Asn	Ser	Ser	Met	Glu	Lys	Lys	Arg	Gln	Glu	Glu
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Glu	Glu	Arg	Ala	Arg	Val	Arg	Arg	Phe	Pro	Ser	Phe	Gln	Pro	Glu	Asp
	50					55					60				

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Phe	Ile	Phe	Met	Asn	Lys	Ala	Glu	Ala	Asn	Asn	Ile	Glu	Lys	Ala	Ala
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Asn	Glu	Ser	Ser	Ser	Ala	Ser	Asn	Glu	Tyr	Asp	Gly	Ser	Lys	Glu	Lys
				85					90					95	
Gln	Glu	Gly	Ser	Glu	Thr	Asn	Val	Cys	Leu	Asp	Leu	Asn	Leu	Ser	Leu
			100					105					110		